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<!--StartFragment-->RESULT 2
PCT-US02-05064-286
; Sequence 286, Application PC/TUS0205064
 GENERAL INFORMATION:
  APPLICANT: Human Genome Sciences, Inc.
  TITLE OF INVENTION: 83 Human Secreted Proteins
  FILE REFERENCE: PS735PCT
  CURRENT APPLICATION NUMBER: PCT/US02/05064
  CURRENT FILING DATE: 2002-03-28
  PRIOR APPLICATION NUMBER: 60/270,658
  PRIOR FILING DATE: 2001-02-23
  PRIOR APPLICATION NUMBER: 60/304,444
  PRIOR FILING DATE: 2001-07-12
  NUMBER OF SEQ ID NOS: 445
  SOFTWARE: PatentIn Ver. 2.0
 SEQ ID NO 286
   LENGTH: 187
   TYPE: PRT
   ORGANISM: Homo sapiens
PCT-US02-05064-286
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 Best Local Similarity 100.0%; Pred. No. 1.8e-98;
 Matches 187; Conservative
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Qу
             Db
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RESULT 3
US-10-644-807-203
; Sequence 203, Application US/10644807
; GENERAL INFORMATION:
  APPLICANT: Rosen et al.
  TITLE OF INVENTION: 83 Human Secreted Proteins
  FILE REFERENCE: PS735
  CURRENT APPLICATION NUMBER: US/10/644,807
  CURRENT FILING DATE: 2003-08-21
  PRIOR APPLICATION NUMBER: PCT/US02/05064
  PRIOR FILING DATE: 2002-02-21
  PRIOR APPLICATION NUMBER: 60/270,658
  PRIOR FILING DATE: 2001-02-23
  PRIOR APPLICATION NUMBER: 60/304,444
  PRIOR FILING DATE: 2001-07-12
  NUMBER OF SEQ ID NOS: 445
  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 203
   LENGTH: 187
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TYPE: PRT
   ORGANISM: Homo sapiens
US-10-644-807-203
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  Best Local Similarity
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                              Pred. No. 1.8e-98;
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             Db
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RESULT 4
US-10-644-807-286
 Sequence 286, Application US/10644807
 GENERAL INFORMATION:
   APPLICANT: Rosen et al.
   TITLE OF INVENTION: 83 Human Secreted Proteins
   FILE REFERENCE: PS735
   CURRENT APPLICATION NUMBER: US/10/644,807
   CURRENT FILING DATE: 2003-08-21
   PRIOR APPLICATION NUMBER: PCT/US02/05064
   PRIOR FILING DATE: 2002-02-21
   PRIOR APPLICATION NUMBER: 60/270,658
   PRIOR FILING DATE: 2001-02-23
   PRIOR APPLICATION NUMBER: 60/304,444
   PRIOR FILING DATE: 2001-07-12
   NUMBER OF SEQ ID NOS: 445
   SOFTWARE: PatentIn Ver. 2.0
  SEO ID NO 286
   LENGTH: 187
    TYPE: PRT
    ORGANISM: Homo sapiens
US-10-644-807-286
  Query Match
                       100.0%;
                              Score 983; DB 36;
                                                Length 187;
  Best Local Similarity
                       100.0%;
                               Pred. No. 1.8e-98;
  Matches 187; Conservative
                             0; Mismatches
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Qу
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Db 121 SQICELKYEKTLDLASVDLRKMRVAELKQILHSWGEECRACAEKTDYVNLIQELAPKYAA 180

Qy 181 THPKTEL 187

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Db 181 THPKTEL 187

<!--EndFragment-->